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1647

RAW SEQUENCE LISTING

DATE: 04/10/2001

PATENT APPLICATION: US/09/187,693

TIME: 12:27:20

Input Set : A:\C420CIP2SL.TXT

Output Set: N:\CRF3\04102001\I187693.raw

ENTERED

4 <110> APPLICANT: Jakobovits, Aya  
5 Yang, Xiao-Dong  
6 Gallo, Michael  
7 Jia, Xiao-Chi  
9 <120> TITLE OF INVENTION: Human Monoclonal Antibodies to Epidermal  
10 Growth Factor Receptor  
13 <130> FILE REFERENCE: Cell 4.20 CIP2  
15 <140> CURRENT APPLICATION NUMBER: 09/187,693  
16 <141> CURRENT FILING DATE: 1998-11-05  
18 <150> PRIOR APPLICATION NUMBER: 09/162,280  
19 <151> PRIOR FILING DATE: 1998-09-29  
21 <150> PRIOR APPLICATION NUMBER: 08/851,362  
22 <151> PRIOR FILING DATE: 1997-05-05  
24 <160> NUMBER OF SEQ ID NOS: 75  
26 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
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29 <211> LENGTH: 22  
30 <212> TYPE: DNA  
31 <213> ORGANISM: human  
33 <400> SEQUENCE: 1  
34 cagggtgcagc tggagcagtc gg 22  
36 <210> SEQ ID NO: 2  
37 <211> LENGTH: 24  
38 <212> TYPE: DNA  
39 <213> ORGANISM: human  
41 <400> SEQUENCE: 2  
42 gctgagggag tagagtcctg agga 24  
44 <210> SEQ ID NO: 3  
45 <211> LENGTH: 294  
46 <212> TYPE: DNA  
47 <213> ORGANISM: human  
49 <400> SEQUENCE: 3  
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51 gggaagggcc tggactgcat tgggtacatc tattacagtg ggagcaccta ctacaaccg 120  
52 tccctcaaga gtcgagttac catatcagta gacacgtcta agaatacagtt ctctctgaag 180  
53 ctgacctctg tgactgccgc ggacacggcc gtgtattact gtgcgagatc tacggttgta 240  
54 aatccggggt ggttcgacct ctggggccar ggaaccctgg tcaccgtctc ctca 294  
56 <210> SEQ ID NO: 4  
57 <211> LENGTH: 264  
58 <212> TYPE: DNA  
59 <213> ORGANISM: human  
61 <400> SEQUENCE: 4  
62 accatcactt gccaggcgag tcaggacatt aacaactatt taaattggtt tcagcagaaa 60  
63 ccagggaag cccctaaggt cctgatccac gatgcatcca atttggaag agggggccca 120  
64 tcaaggttca gtggaagtgg atctgggaca gattttactt tcaccatcag cggcctgcag 180  
65 cctgaagaca ttgcaacata ttattgtcaa cagtatgaaa gtctcccact cactttcggc 240  
66 ggagggacca aggttgagat caaa 264

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68 <210> SEQ ID NO: 5
69 <211> LENGTH: 291
70 <212> TYPE: DNA
71 <213> ORGANISM: human
73 <400> SEQUENCE: 5
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75 ggaagggcc tggagtggat tgggtccatc tattacagtg ggaacacctt ctacaaccgc      120
76 tccctcaaga gtcgagttac catatcacta gacacgtcta agaaccagtt ctccctgaag      180
77 ctgagttctg tgactgccgc ggacacggcc gtgtgttact gtgcgagaaa tatagtgact      240
78 acgggtgctt ttgatattctg gggccaaggg acaatggtea ccgtctcttc a          291
80 <210> SEQ ID NO: 6
81 <211> LENGTH: 264
82 <212> TYPE: DNA
83 <213> ORGANISM: human
85 <400> SEQUENCE: 6
86 accatcactt gtcaggcgag tcaggacatt accatttatt taaattggta tcaacagaaa      60
87 ccagggaaaag ccctaagct cctgatcaac gacgcattca gtttggaaac aggggtccca      120
88 ttaaggttca gtggaagtgg atctgggaca gattttactt tcaccatcag cagcctgcag      180
89 cctgaagata ttgaacata ttactgtcaa cagtatgata atctccgcgt cactttcggc      240
90 ggcgggacca aggtggcgat caaa          264
92 <210> SEQ ID NO: 7
93 <211> LENGTH: 288
94 <212> TYPE: DNA
95 <213> ORGANISM: human
97 <400> SEQUENCE: 7
98 gtctctggtg gctccatcag cagtggatgat tactactgga cctggatccg ccagcaccga      60
99 ggaagggcc tggagtggat tgggtacatc tattacagtg ggaacacctt ctacaaccgc      120
100 tccctcaaga gtcgagtttc catgtcaata gacacgtctg agaaccagtt ctccctgaag      180
101 ctgagctctg tgactgccgc ggacacggcc gtgtattact gtgcgagaaa accagtgact      240
102 gggggggagg actactgggg ccagggaaacc ctggtcaccg tctcctca          288
104 <210> SEQ ID NO: 8
105 <211> LENGTH: 262
106 <212> TYPE: DNA
107 <213> ORGANISM: human
109 <400> SEQUENCE: 8
110 accatcactt gccaggcgag tcaggacatt agtaactatt taaattggta tcagcagaaa      60
111 ccagggaaaag ccctaagctc ctgatctacg atgcttccaa tttggaaaca ggggtcccat      120
112 caaggttcag tggagtggat ctgggacaga ttttactttc accatcagca gcctgcagcc      180
113 tgaagatgtt ggaacatatg tctgtcaaca gtatgagagt ctcccgtagc gttttggcca      240
114 ggggacaaaa ctggagatca aa          262
116 <210> SEQ ID NO: 9
117 <211> LENGTH: 291
118 <212> TYPE: DNA
119 <213> ORGANISM: human
121 <220> FEATURE:
122 <221> NAME/KEY: misc_feature
123 <222> LOCATION: (1)...(291)
124 <223> OTHER INFORMATION: n = A,T,C or G
126 <400> SEQUENCE: 9

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	127	gtctctggtg	gctccatcaa	cagtgggtgat	ttctactgga	gctggatccg	ccaacaccca	60
	128	gggaagggcc	tggagtggat	tgggtacatc	tattacagt	ggagcaccta	ctacaacccg	120
	129	tccctcaaga	gtcgagttac	catgtcaata	gaccgcgtcta	agaaccagtt	ctccctgaaa	180
W-->	130	ctgatctctg	tgactgccgc	ggacacggcc	gtttattact	gtgcgacntc	cctttactat	240
	131	ggcgggggta	tggacgtctg	gggccaaagg	accacggtca	cgtctcctc	a	291
	133	<210>	SEQ ID NO: 10					
	134	<211>	LENGTH: 264					
	135	<212>	TYPE: DNA					
	136	<213>	ORGANISM: human					
	138	<220>	FEATURE:					
	139	<221>	NAME/KEY: misc_feature					
	140	<222>	LOCATION: (1)...(264)					
	141	<223>	OTHER INFORMATION: n = A,T,C or G					
	143	<400>	SEQUENCE: 10					
	144	accatcactt	gccaggcgag	tcaggacatt	aacaactatt	tgaattggta	tcagcagagg	60
W-->	145	cnngggaacg	cccctaaact	cctgatctac	gatgcatcca	atttggaaac	aggggtccca	120
	146	tcaaggttca	gtggaagtgg	atctgggaca	gattttactt	tcaccatcaa	cagcctgcag	180
	147	cctgaagata	ttgcgacata	ttattgtcaa	cactatgatc	atctcccgtg	gacgttcggc	240
W-->	148	caagggacca	aggtggaant	caaa				264
	150	<210>	SEQ ID NO: 11					
	151	<211>	LENGTH: 291					
	152	<212>	TYPE: DNA					
	153	<213>	ORGANISM: human					
	155	<400>	SEQUENCE: 11					
	156	gtctctggtg	gctccatcaa	caatgggtgat	tactactgga	gctggatccg	ccagcaccca	60
	157	gggaagggcc	tggagtggat	tgggcacatc	tattacagt	ggagcaccta	ctacatcccg	120
	158	tccctcaaga	gtcgaactac	catatcagta	gacacgtcta	agaaccagtt	ctccctgaag	180
	159	ctgaactctg	tgactgccgc	ggacacggcc	gtgtattact	gtgcgagagg	gacagtaact	240
	160	acgtactact	ttgactactg	gggccagggg	accctgggtca	cgtctcctc	a	291
	162	<210>	SEQ ID NO: 12					
	163	<211>	LENGTH: 270					
	164	<212>	TYPE: DNA					
	165	<213>	ORGANISM: human					
	167	<400>	SEQUENCE: 12					
	168	accatcactt	gccgggcaag	tcagagcatt	agcagctatt	taaattggta	tcagcagaaa	60
	169	ccagggaaa	cccctaagct	cctgatctat	gctgcatcca	gtttgcaaag	tgggggtccca	120
	170	tcaaggttca	gtggcagtgg	atctgggaca	gatttcactc	tcaccatcag	cagtctgcaa	180
	171	cctgaagatt	ttgcaactta	ctactgtcaa	cagggttaca	gaaccctctc	ggagtgcagt	240
	172	tttggccagg	ggaccaagct	ggagatcaaa				270
	174	<210>	SEQ ID NO: 13					
	175	<211>	LENGTH: 291					
	176	<212>	TYPE: DNA					
	177	<213>	ORGANISM: human					
	179	<400>	SEQUENCE: 13					
	180	gtctctggtg	gctccgtcag	cagtgggtgat	tactactgga	gctggatccg	gcagccccc	60
	181	gggaagggac	tggagtggat	tggacatctc	tattacagt	ggaacaccaa	ctacaacccc	120
	182	tccctcaaga	gtcgagtcac	catatcatta	gacacgtcca	agaaccagtt	ctccctgaag	180
	183	ctgagctctg	tgaccgtctg	ggacacggcc	gtgtattact	gtgcgagata	ttttttgact	240
	184	qattccttct	ttgactactg	gggccagggg	accctgattca	cgtctcctc	a	291

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186 <210> SEQ ID NO: 14
187 <211> LENGTH: 264
188 <212> TYPE: DNA
189 <213> ORGANISM: human
191 <400> SEQUENCE: 14
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193 ccaggggaaag cccctaagct cctgatcaac gatgcatccg atttggaaac aggggtccca      120
194 tcaaggatca gtggaagtgg atctgggaca gattttactt tcaccatcag caacctgcag      180
195 cctgaagata ttgcaacata ttactgtcaa caatatgata gtctcccgct cactttcggc      240
196 ggagggacca aggtggagat caga                                     264
198 <210> SEQ ID NO: 15
199 <211> LENGTH: 288
200 <212> TYPE: DNA
201 <213> ORGANISM: human
203 <400> SEQUENCE: 15
204 gtctctggtg gctccgtcta cagtggatgat tactactgga gctggatccg gcagccccc      60
205 gggaagggac tggagtggat tgggtatata tattacagtg ggagcaccaa ttacaatccc      120
206 tccctcaaga gtcgagtcac catatcagta gacacgtcca agaaccagtt ctccctgaag      180
207 ctgagctctg tgaccgctgc ggacacggcc gtgtattact gtgcgagaga ctccatactg      240
208 ggagctacca actactgggg ccaggggaacc ctggtcaccg tctcctca      288
210 <210> SEQ ID NO: 16
211 <211> LENGTH: 264
212 <212> TYPE: DNA
213 <213> ORGANISM: human
215 <220> FEATURE:
216 <221> NAME/KEY: misc_feature
217 <222> LOCATION: (1)...(264)
218 <223> OTHER INFORMATION: n = A,T,C or G
220 <400> SEQUENCE: 16
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222 ccaggggaaag cccctaaast cctgatctcc gatgcatcca atttagaaac aggggtccca      120
W--> 223 tcgaggttca gtggaagtgg atctgggaca gantntactt tcaccatcag cagcctgcag      180
W--> 224 cctgaagata ttgcnacata tcactgtcna cagtatnata gtctcccgct cactttcggc      240
225 ggagggacca aggtagagat caaa                                     264
227 <210> SEQ ID NO: 17
228 <211> LENGTH: 288
229 <212> TYPE: DNA
230 <213> ORGANISM: human
232 <400> SEQUENCE: 17
233 gtctctggtg gctccgtcag cagtggatgat tactactgga cctggatccg gcagtcccca      60
234 gggaagggac tggagtggat tggacacatc tattacagtg ggaacaccaa ttataacccc      120
235 tccctcaaga gtcgactcac catatcaatt gacacgtcca agactcagtt ctccctgaag      180
236 ctgagttctg tgaccgctgc ggacacggcc atttattact gtgtgcgaga tcgagtgact      240
237 ggtgcttttg atatctgggg ccaagggaca atggtcaccg tctcttca      288
239 <210> SEQ ID NO: 18
240 <211> LENGTH: 264
241 <212> TYPE: DNA
242 <213> ORGANISM: human
244 <400> SEQUENCE: 18

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245 accatcactt gccaggcgag tcaggacatc agcaactatt taaattggta tcagcagaaa      60
246 ccaggggaaag cccctaaact cctgatctac gatgcatcca atttggaaac aggggtccca      120
247 tcaaggttca gtggaagtgg atctgggaca gattttactt tcaccatcag cagcctgcag      180
248 cctgaagata ttgcaacata tttctgtcaa cactttgatc atctcccgct cgctttcggc      240
249 ggagggacca aggtggagat caaa                                     264
251 <210> SEQ ID NO: 19
252 <211> LENGTH: 480
253 <212> TYPE: DNA
254 <213> ORGANISM: human
256 <220> FEATURE:
257 <221> NAME/KEY: misc_feature
258 <222> LOCATION: (1)...(480)
259 <223> OTHER INFORMATION: n = A,T,C or G
261 <400> SEQUENCE: 19
W--> 262 gcgatccagc cttttaggtc catgccnttc tctgtgnag cgtctggatt ccccttcagt      60
W--> 263 agntntggca tgcactgggt ccgccaggct ccaggcaagg ggctggagtg ggtggcagtt      120
264 atatggtatg atggaagtaa taaatactat gcagactocg tgaagggccg attcaccatc      180
265 tccagagaca attccaagaa cacgctgtat ctgcaaatga acagactgag agccgaggac      240
266 acggctgtgt attactgtgc gagatttttg gagtggttac ctttgacta ctggggccag      300
W--> 267 ggaaccctgg tcaccgtctn ctcagactcc accaagggcc catcgggtctt cncctggcg      360
W--> 268 ccctgcttcc aggagcacc ctcnganagca canangggcc ctgggactgn ctgntacaag      420
W--> 269 gactnctttc cctcnaaccn ggtgaccntn tcntgggaaa ctcagngcnc ntctnnatna      480
271 <210> SEQ ID NO: 20
272 <211> LENGTH: 490
273 <212> TYPE: DNA
274 <213> ORGANISM: human
276 <220> FEATURE:
277 <221> NAME/KEY: misc_feature
278 <222> LOCATION: (1)...(490)
279 <223> OTHER INFORMATION: n = A,T,C or G
281 <400> SEQUENCE: 20
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283 gggcattagc aatttttttag cctggtttca gcagaaacca gggatagccc ctaagtccct      120
284 gatctatgct gcatccactt tgcaaaagg ggtcccatca aagttcaccc gcagtggata      180
285 tgggacagat ttcactctca ccatcagcag cctgcagcct gaagactttg caacttatta      240
286 ttgtcaacaa tataatgttt acccatcac tttcggccct gggaccaaag tggatatcaa      300
287 acgaactgtg gctgcacat ctgtcttcac cttccgccca tctgatgagc aagttgaaat      360
288 ctggaactgc ctctgtttg tgctgtctga ataacttcta tcccagagag gccaaagtac      420
W--> 289 agtggagggt ggataacgcc ncnnttggeg gnntcctttc nctcncct cctcnnccn      480
W--> 290 cctctcncna                                     490
292 <210> SEQ ID NO: 21
293 <211> LENGTH: 449
294 <212> TYPE: DNA
295 <213> ORGANISM: human
297 <220> FEATURE:
298 <221> NAME/KEY: misc_feature
299 <222> LOCATION: (1)...(449)
300 <223> OTHER INFORMATION: n = A,T,C or G
302 <400> SEQUENCE: 21

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FYI

**Please Note:**

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a c rresp nding explanation is presented in the <220> to <223> fields of each sequence which pr sents at least one n or Xaa.

## VERIFICATION SUMMARY

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Input Set : A:\C420CIP2SL.TXT

Output Set: N:\CRF3\04102001\I187693.raw

L:130 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9  
L:145 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10  
L:148 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10  
L:221 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16  
L:223 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16  
L:224 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16  
L:262 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19  
L:263 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19  
L:267 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19  
L:268 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19  
L:269 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19  
L:282 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20  
L:289 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20  
L:290 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20  
L:308 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21  
L:309 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21  
L:310 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21  
L:323 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22  
L:324 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22  
L:329 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22  
L:343 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23  
L:350 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23  
L:364 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24  
L:391 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25  
L:411 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26  
L:424 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27  
L:425 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27  
L:426 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27  
L:429 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27  
L:430 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27  
L:443 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28  
L:449 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28  
L:450 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28  
L:463 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29  
L:464 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29  
L:465 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29  
L:466 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29  
L:467 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29  
L:468 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29  
L:481 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30  
L:482 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30  
L:487 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30  
L:488 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30  
L:501 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31  
L:506 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31  
L:507 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31  
L:508 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31  
L:521 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32

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L:528 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32

L:557 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34